In re Application of:
PATENT
Frost et al.
Attorney Docket No. HALO1330-1

Application No.: 10/539,110 Filing Date: April 19, 2006

Page 2

AMENDMENTS TO THE CLAIMS

Please amend claims 1, 50 and 58 as set forth below.

Please amend and rejoin claims 40-56 (see proposed amendments upon rejoinder).

Please cancel claim 59 without prejudice.

The current listing of claims replaces all prior listings.

Listing of Claims

- (Currently Amended) A substantially purified <u>soluble</u> mammalian chondroitinase glycoprotein (CHASEGP) comprising a <u>soluble HYAL4 polypeptide encoded by a nucleic acid</u> sequence as set forth in SEO ID NO:5-6.
- 2. (Withdrawn) The glycoprotein of claim-1, wherein the polypeptide is selected from the group of a polypeptide that comprises a sequence of amino acids encoded by nucleotides 642-2087 in SEQ ID No. 3 and includes at least about 74% amino acid sequence identity with the sequence of amino acids set forth in SEQ ID No. 1; a polypeptide that comprises a sequence of amino acids encoded by the sequence of nucleotides set forth in SEQ ID No. 2; a polypeptide that comprises a sequence of amino acids encoded by a sequence of nucleotides that hybridizes along at least 85% of its full-length under conditions of high stringency to the sequence of nucleotides set forth as nucleotides 642-2087 in SEQ ID No. 3.
- (Previously Presented) The glycoprotein of claim 1, wherein a sugar moiety is covalently
 attached to an asparagine residue selected from the group consisting of amino acid residues 52,
 81 and 309 as set forth in SEO ID NO:6.
- (Previously Presented) The glycoprotein of claim 3, wherein said sugar moiety is covalently linked to said glycoprotein through a asparagine peptide specific N-glycosidase F (PNGase) sensitive bond.
- (Previously Presented) The glycoprotein of claim 3, wherein said sugar moiety is of the high mannose type.

In re Application of:

PATENT
Frost et al.

Attornev Docket No. HALO1330-1

Application No.: 10/539,110 Filing Date: April 19, 2006

Page 3

(Previously Presented) The glycoprotein of claim 3, wherein said sugar moiety is of the complex type.

- (Previously Presented) The glycoprotein of claim 3, wherein said sugar moiety is of the hybrid type.
- 8. (Previously Presented) The glycoprotein of claim 3, wherein said sugar moiety is substantially terminated with sialic acid.

(Canceled)

- 10. (Withdrawn) The substantially purified glycoprotein of claim 1, wherein the chondroitinase domain comprised the sequence of amino acids set forth as amino acids 35-457 of SEQ ID No.1.
- 11. (Withdrawn) The substantially purified glycoprotein of claim 1 that has more that about 80% sequence identity with a polypeptide that comprises the sequence of amino acids set forth as SEQ ID No. 1 or as the sequence of amino acids set forth as SEQ ID No. 2, wherein the polypeptide is a chondroitinase.
- 12. (Previously Presented) The glycoprotein of claim 1, wherein a chondroitinase domain or the catalytically active portion thereof is encoded by a nucleic acid molecule that hybridizes under conditions of high stringency along at least 70% of its full-length to a nucleic acid molecule comprising a sequence of nucleotides set forth as nucleotides 726-1995 in SEQ ID No. 3 or as set forth in SEQ ID. No. 5.
- 13. (Previously Presented) The glycoprotein of claim 1, wherein the CHASEGP is a human polypeptide.
- 14. (Previously Presented) The glycoprotein of claim 1, wherein the polypeptide is encoded by a nucleic acid sequence as set forth in SEQ ID NO. 5.
- (Canceled)

In re Application of:
PATENT
Frost et al.
Attorney Docket No. HALO1330-1

Application No.: 10/539,110 Filing Date: April 19, 2006

Page 4

16. (Previously Presented) The glycoprotein of claim 1, wherein the polypeptide further comprises one or more domains selected from a polyhistidine domain, protein A domain, FLAG domain, Factor XA domain, or an eneterokinase domain, or a combination thereof.

- 17. (Withdrawn) A glycoprotein of claim 1 that is a mutein, wherein: up to about 50% of the amino acids are replaced with another amino acid; and the resulting polypeptide is a single chain or two chain polypeptide that has catalytic activity of at least 10% of the unmutated polypeptide.
- 18. (Withdrawn) The glycoprotein of claim 17, wherein up to about 10% of the amino acids are replaced with another amino acid.
- 19. (Withdrawn) The glycoprotein of claim 17, wherein the resulting polypeptide is a single chain or two chain polypeptide and has catalytic activity of at least 50% of the unmutated polypeptide.
- 20. (Withdrawn) The glycoprotein of claim 17, wherein a free Cysteine in the chondroitinase domain is replaced with another amino acid
- 21. (Withdrawn) The glycoprotein of claim 20, wherein the replacing amino acid is a serine.
- (Canceled)
- 23. (Withdrawn) A nucleic acid molecule, comprising a sequence of nucleotides that encodes the polypeptide of claim 1.
- 24. (Withdrawn) The nucleic acid molecule of claim 23 that comprises a sequence of nucleotides selected from the group consisting of: (a) a sequence of nucleotides set forth as nucleotides 726-1995 in SEQ ID No. 3; (b) a sequence of nucleotides that hybridizes under high stringency along its length or along at least about 70% of the full-length to the sequence of nucleotides set forth as nucleotides 726-1995 in SEQ ID No. 3 or as SEQ ID No. 5 (c) a sequence of nucleotides that encodes the polypeptide of SEQ ID No. 6; (d) a sequence of nucleotides that is a splice variant of a, b, or c); (e) a sequence of nucleotides that encodes the

In re Application of:

Frost et al.

Attorney Docket No. HALO1330-1

Application No.: 10/539,110 Filing Date: April 19, 2006

Page 5

chondroitinase domain or a catalytically active portion thereof that includes a sequence of nucleotides having at least about 60%, 70%, 80%, 90% or 95% sequence identity the sequence set forth in SEQ ID Nos. 3,4 or 5; and (f) a sequence of nucleotides comprising degenerate codons of (a), (b),(c), (d) or (e).

- 25. (Withdrawn) An isolated nucleic molecule that encodes a mutein of claim 17.
- 26. (Withdrawn) A vector comprising the nucleic acid molecule of claim 23.
- 27. (Withdrawn) The vector of claim 26 that is an expression vector.
- 28. (Withdrawn) The vector of claim 26 that is a eukaryotic vector.
- (Withdrawn) The vector of claim 26 that includes a sequence of nucleotides that directs secretion of any polypeptide encoded by a sequence of nucleotides operatively linked thereto.
- 30. (Withdrawn) The vector of claim 26 that is a Pichia vector or an E. coli vector.
- 31. (Withdrawn) A cell, comprising the vector of claim 26.
- 32. (Withdrawn) The cell of claim 31 that is a prokaryotic cell.
- 33. (Withdrawn) The cell of claim 31 that is a eukaryotic cell.
- 34. (Withdrawn) The cell of claim 31 that is selected from among a bacterial cell, a yeast cell, a plant cell, an insect cell and an animal cell.
- 35. (Withdrawn) The cell of claim 31 that is a mammalian cell.
- 36. (Withdrawn) A nucleic acid molecule encoding a polypeptide of claim 1.
- 37. (Withdrawn) A vector, comprising nucleic acid molecule of claim 23.
- 38. (Withdrawn) A cell, comprising the vector of claim 23.

Application No.: 10/539,110 Filing Date: April 19, 2006

Page 6

- 39. (Withdrawn) A recombinant non-human animal, wherein an endogenous gene that encodes a polypeptide of claim 1 has been deleted or inactivated by homologous recombination or insertional mutagenesis of the animal or an ancestor thereof.
- 40. (Withdrawn) A method for generating soluble recombinant CHASEGP comprising, introduction of a nucleic acid as described in SEQ ID NO: 4 set forth in SEQ ID NO:5 operably linked to a suitable promoter into a eukaryotic cell capable of incorporating said N-linked sugar moieties into CHASEGP.
- 41. (Withdrawn) The method of claim 40, wherein the eukaryotic cell is mammalian.
- 42. (Withdrawn) The method of claim 40, wherein said eukaryotic cell is an insect.
- 43. (Withdrawn) The method of claim 40, wherein said eukaryotic cell is a yeast
- 44. (Withdrawn) The method of claim 3, wherein said eukaryotic cell is a plant.
- 45. (Withdrawn) The method of claim 40, wherein the expressible polynucleotide is introduced into a cell ex vivo, thereby generating a genetically modified cell containing the expressible polynucleotide, and wherein administering the expressible polynucleotide to the subject comprises administering the genetically modified cell to the subject.
- 46. (Withdrawn) The method of claim 45, wherein the cell is autologous with respect to the subject.
- (Withdrawn) The method of claim 45, wherein the cell is haplotype matched with respect to the subject.
- 48. (Withdrawn) A method for generating the CHASEGP a soluble glycoprotein comprising, contacting a chondroitinase- the polypeptide of claim 1 with glycosyltransferase enzymes capable of introducing said N-linked sugar moieties to generate CHASEGP the polypeptide of claim 1.

Application No.: 10/539,110 Filing Date: April 19, 2006

Page 7

49. (Withdrawn) The method of claim 48 wherein the glycosyltransferase enzymes are derived from canine microsomal membranes

- (Currently amended) A composition comprising a substantially purified <u>soluble</u> chondroitinase glycoprotein (CHASEGP) in a suitable pharmaceutical carrier, wherein the CHASEGP is a <u>soluble HYAL4 polypeptide encoded by a nucleic acid sequence as</u> set forth in SEO ID NO:5-6.
- 51. (Withdrawn) A method for treating an animal a subject suffering from an excess of CHASEGP substrate, said method comprising administration of a recombinant CHASEGP administering the glycoprotein of claim 1 to the subject in an amount sufficient to remove said CHASEGP substrate.
- (Withdrawn) The method of claim 51, wherein said excess substrate is produced from a scar tissue.
- 53. (Withdrawn) The method of claim 52, wherein said scar tissue is a glial scar resulting from spinal cord injury.
- 54. (Withdrawn) The method of claim 52, wherein said scar tissue is a result of surgery.
- 55. (Withdrawn) The method of claim 52, wherein said scar is a keloid scar.
- (Withdrawn) The method of claim 51 wherein said substrate is associated with a hemiated disk.
- (Canceled)
- 58. (Currently amended) The composition of claim 50, wherein the glycoprotein is encoded by a nucleic acid sequence as set forth in SEQ ID NO:6-5.
- (Canceled)